

**Table 5 on page 22:**

TABLE 5  
BLAST Results for Sequences Encoding Polypeptides  
Homologous to NPR1

Clone	Status	SEQ ID NO:	BLAST pLog Score
rr1.pk0001.a11:cgs	CGS	16	1773295 100.00

**Table 6 on page 22:**

TABLE 6  
Percent Identity of Amino Acid Sequences Deduced From the Nucleotide Sequences  
of cDNA Clones Encoding Polypeptides Homologous to NPR1

Clone	SEQ ID NO:	Percent Identity to 1773295
cdt1c.pk001.l6	2	45.8
rr1.pk0001.a11	4	47.6
wre1n.pk0122.c2	6	40.0
p0006.cbyvc82rx	8	40.3
rl0n.pk0063.d10: fis	10	42.1
rr1.pk0001.a11: fis	12	39.1
wre1n.pk0122.c2: fis	14	33.7
rr1.pk0001.a11:cgs	16	34.1

**IN THE CLAIMS:**

**Please cancel claims 7-23** without prejudice to or disclaimer of the subject matter recited therein.

**Please amend claims 1, 5-6, and 24-25 as follows** (a marked-up version showing changes made is attached hereto):

1. (Amended) An isolated polynucleotide comprising:
  - (a) a nucleotide sequence encoding a polypeptide having NPR1 activity, wherein the polypeptide has an amino acid sequence of at least 80% sequence identity based on the Clustal method of alignment when compared to a polypeptide selected from the group consisting of SEQ ID NOs:4, 12, and 16, or